

FIG.1

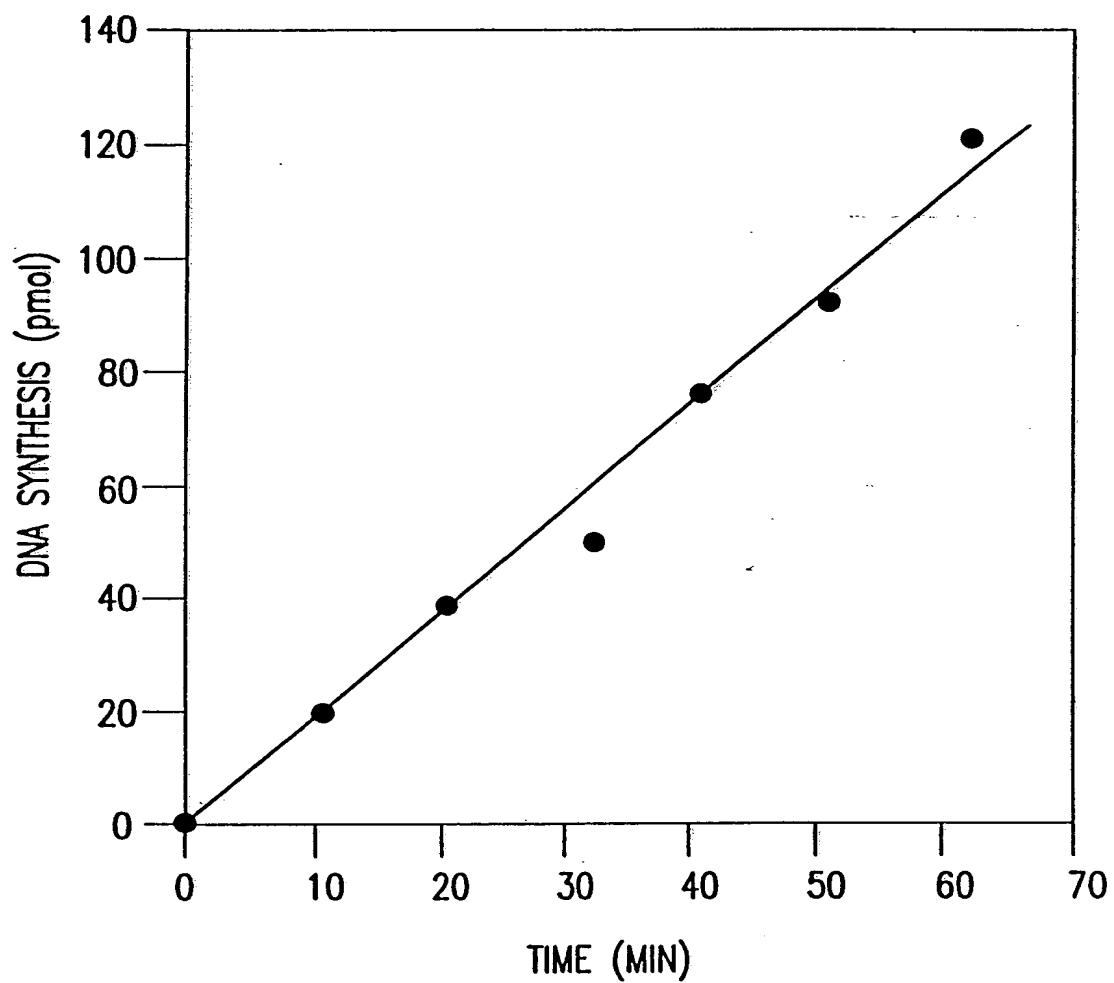


FIG.2

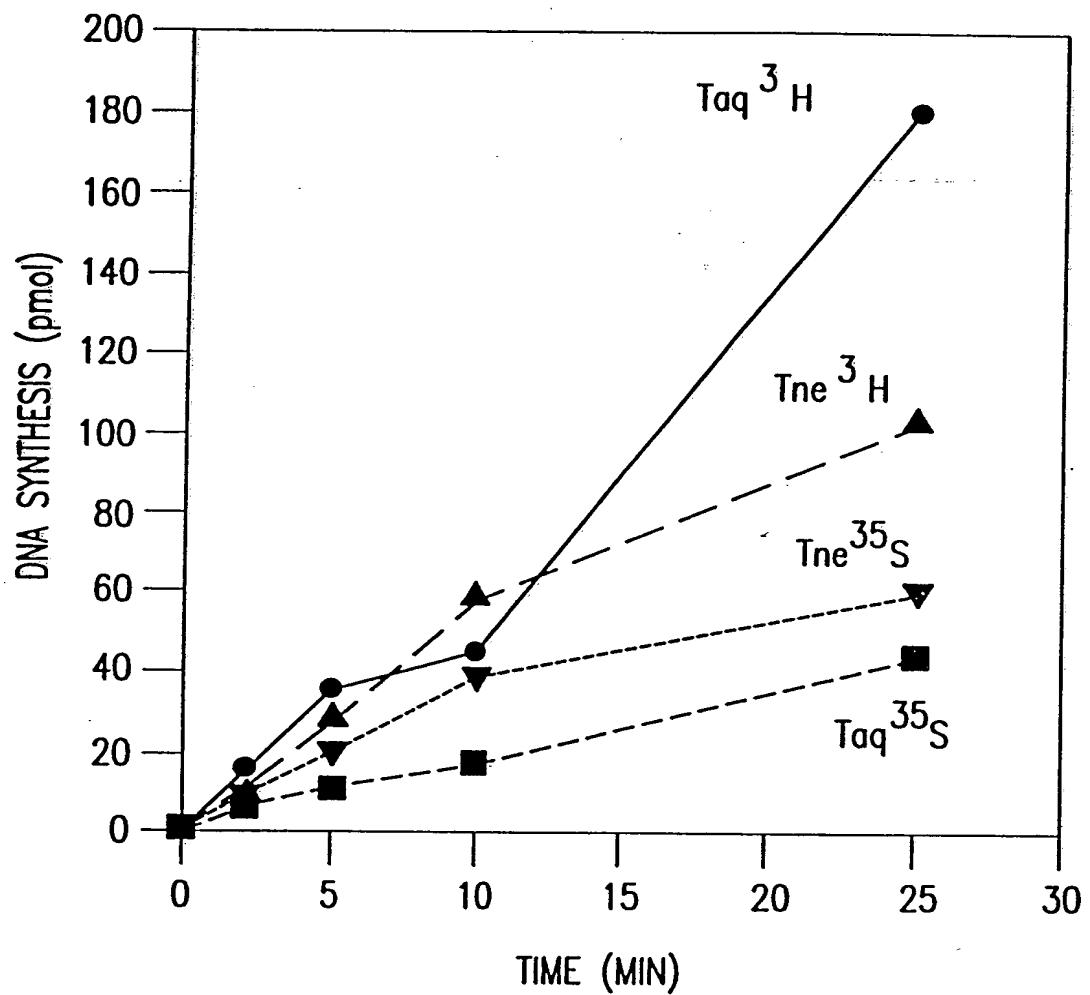


FIG.3

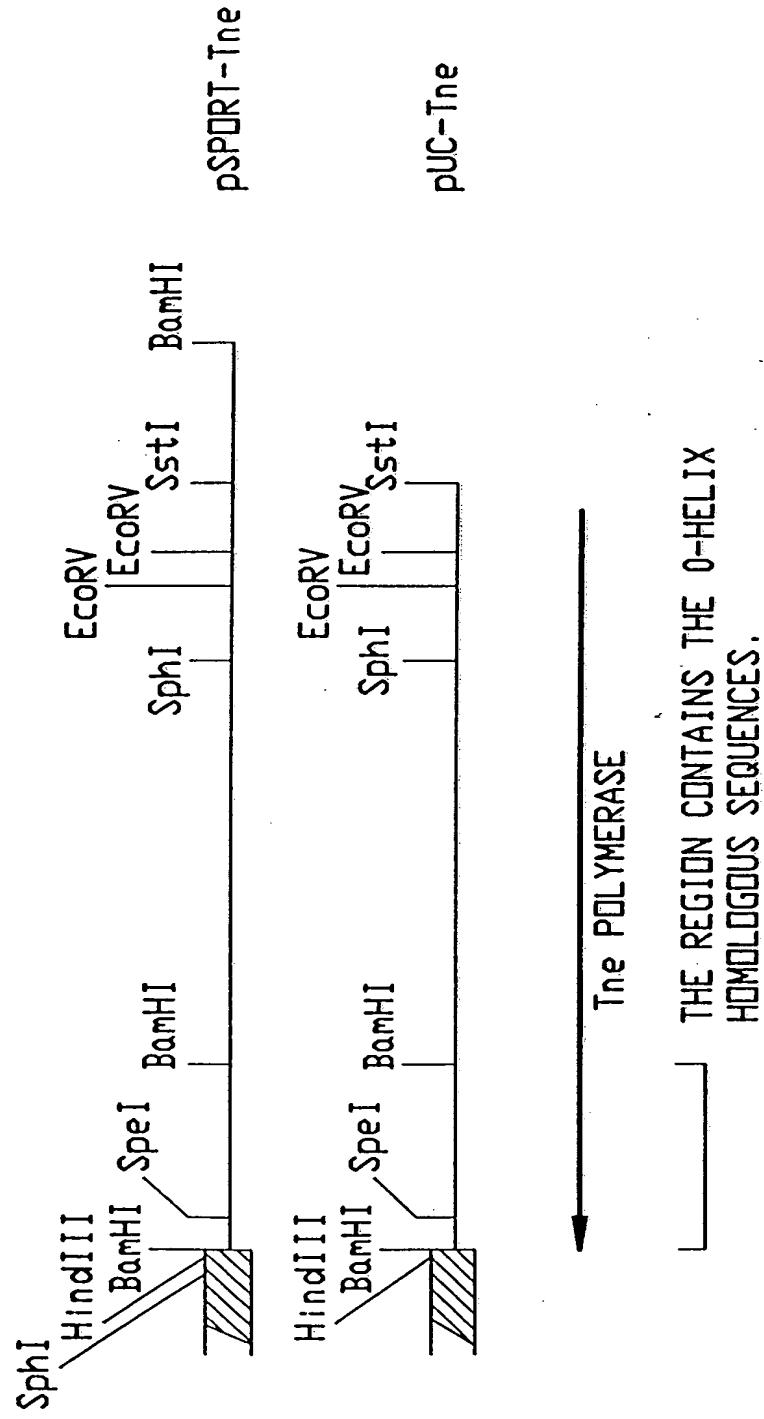


FIG. 4



BamHI

1 CGATCCAGAC TGGTGGATCG TCAGTCGGA TTATTCCCAA ATAGAACTCA GAATCCTCGC  
 G S R L V D R Q C G L F P N R T Q N P R  
 → D P D W W I V S A D Y S Q I E L R I L  
 I Q T G G S S V R I I P K - N S E S S

61 TCATCTCACT CGTGATGAGA ACCTGTGAA GCCCTTCGAG GAGGGCATCG ATGTGCACAC  
 S S Q W - - E P C E G L R G G H R C A H  
 → A H L S G D E N L V K A F E E G I D V H  
 L I S V V M R T L - R P S R R A S M C T

121 CTGACTGCC TCCAGGATCT ACAACGTAAGA CCCAGAAGAA GTGAACGAAG AAATGCCACC  
 L D C L Q D L Q R K A R R S E R R N A T  
 → T L T A S R I Y N V K P E E V N E E M R  
 P - L P P G S T T - S Q K K - T K K C D

181 GGTTGAAAG ATGGTGAAT TCTCTATAAT ATACGGTGTACG ACACCGTACG GTCTTCTGT  
 G W K D G E L L Y N I R C H T V R S F C  
 → R V G K M V N (F) S I I Y G V T P Y G L S  
 G L E R W - T S L - Y T V S H R T V F L

241 GAGACTTCCA ATACCGGTTA AAGAACGAG AAAGATGATT ATCAGCTATT TCACACTGTA  
 E T W N T G - R S R K D D Y Q L F H T V  
 → V R L G I P V K E A E K M I I S Y F T L  
 - D L E Y R L K K Q K R - L S A I S H C

301 TCCAAAGGTG CGAACCTACA TCCAGGAGT TGTTCCAGAG GCAAAAGAGA AGGGCTACGT  
 S K G A K L H P A G C C R G K R E G L R  
 → Y P K V R S Y I Q Q V V A E A K E K G Y  
 I Q R C E A T S S R L L Q R Q K R R A T

361 CAGGACTCTC TTTGGAACAA AAAGAGATAT TCCCCAGCTC ATGGCAAGGG ACAAGAACAC  
 Q D S L W K K K R Y S P A H G K G Q E H  
 → V R T L F G R K R D I P Q L M A R D K N  
 S G L S L E E K E I F P S S W Q G T R T

421 CCAGTCCGAA GCGCAAAGAA TCGCAATAAA CACCCCCATT CAGGAACTG CGGCAGATAT  
 P V R R R K N R N K H P H S G N C G R Y  
 → T Q S E G E R I A I N T P I Q G T A A D  
 P S P K A K E S Q - T P P F R E L R Q I

FIG.5A



481 AATAAAATTG GCTATGATAG ATATAGACGA GGAGCTGAGA AAAAGAAACA TCAAATCCAG  
N K I G Y D R Y R R G A E K K K H E I Q  
→ I I K L A M I D I D E E L R K R N M K S  
-- N W L -- I - T R S - E K E T - N P

541 AATGATCATT CAGGTTCATG ACCAACTGGT CTTGAGGTT CCCGATGAGG AAAAGAAAGA  
N D H S G S - R T G L R G S R - G K R R  
→ R M I I Q V H D E L V F E V P D E E K E  
E - S F R F M T N W S S R F P M R - K K K

601 ACTAGTTGAT CTGGTGAAGA ACAAAATGAC AAATGTGGTG AAAACTCTCTG TGCCTCTTGA  
T S - S G E E Q N D K C G E T L C A S -  
→ E L V D L V K N K M T N V V K L S V P L  
N - L I W - R T K - Q M W - N S L C L L

661 CGTTGACATA ACCATCCGAA AAACCTGGTC TTGA  
G - H K H R K K L V L  
→ E V D I S I G K S W S -  
R L T - A S E K A G L

FIG.5B

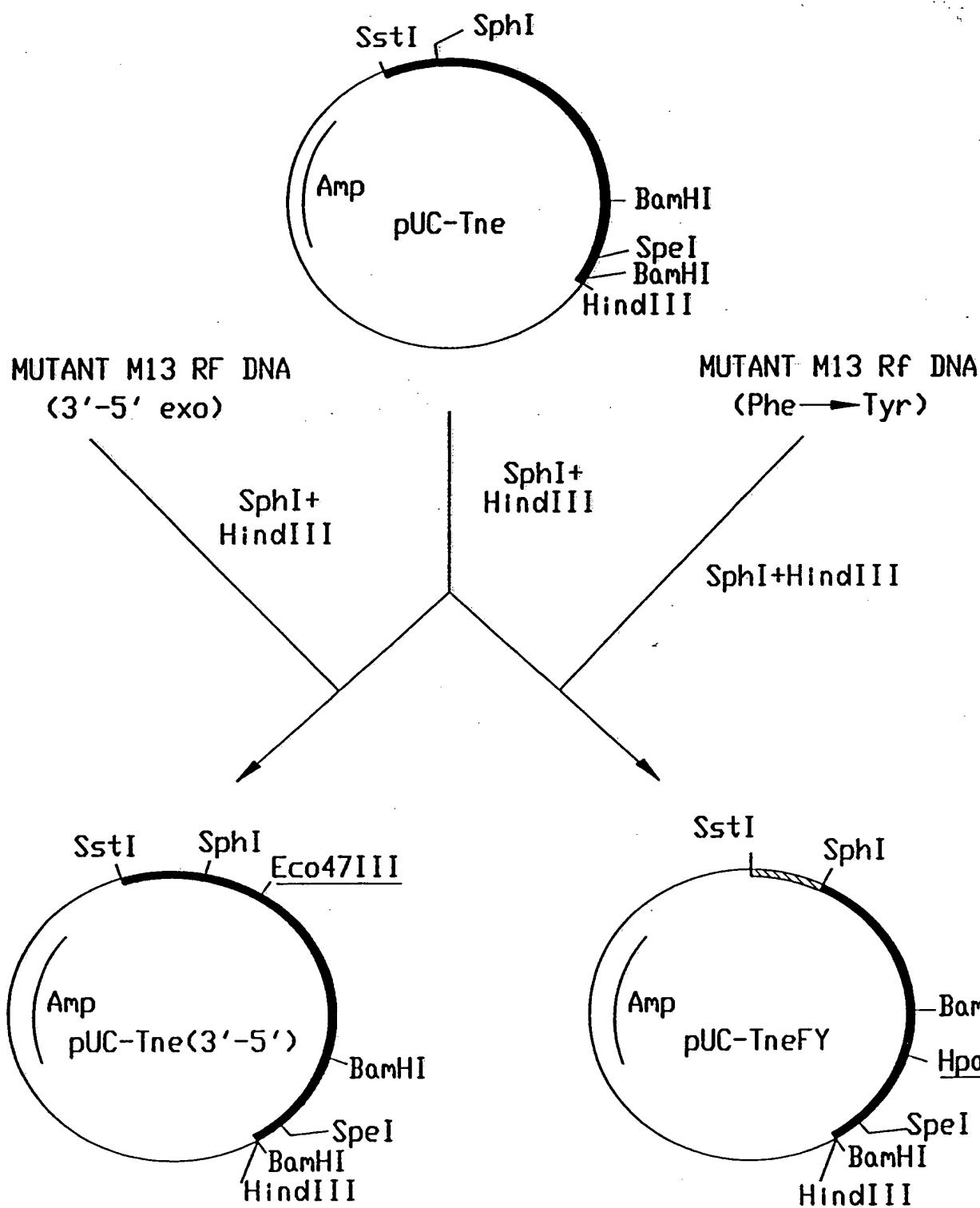


FIG.6A

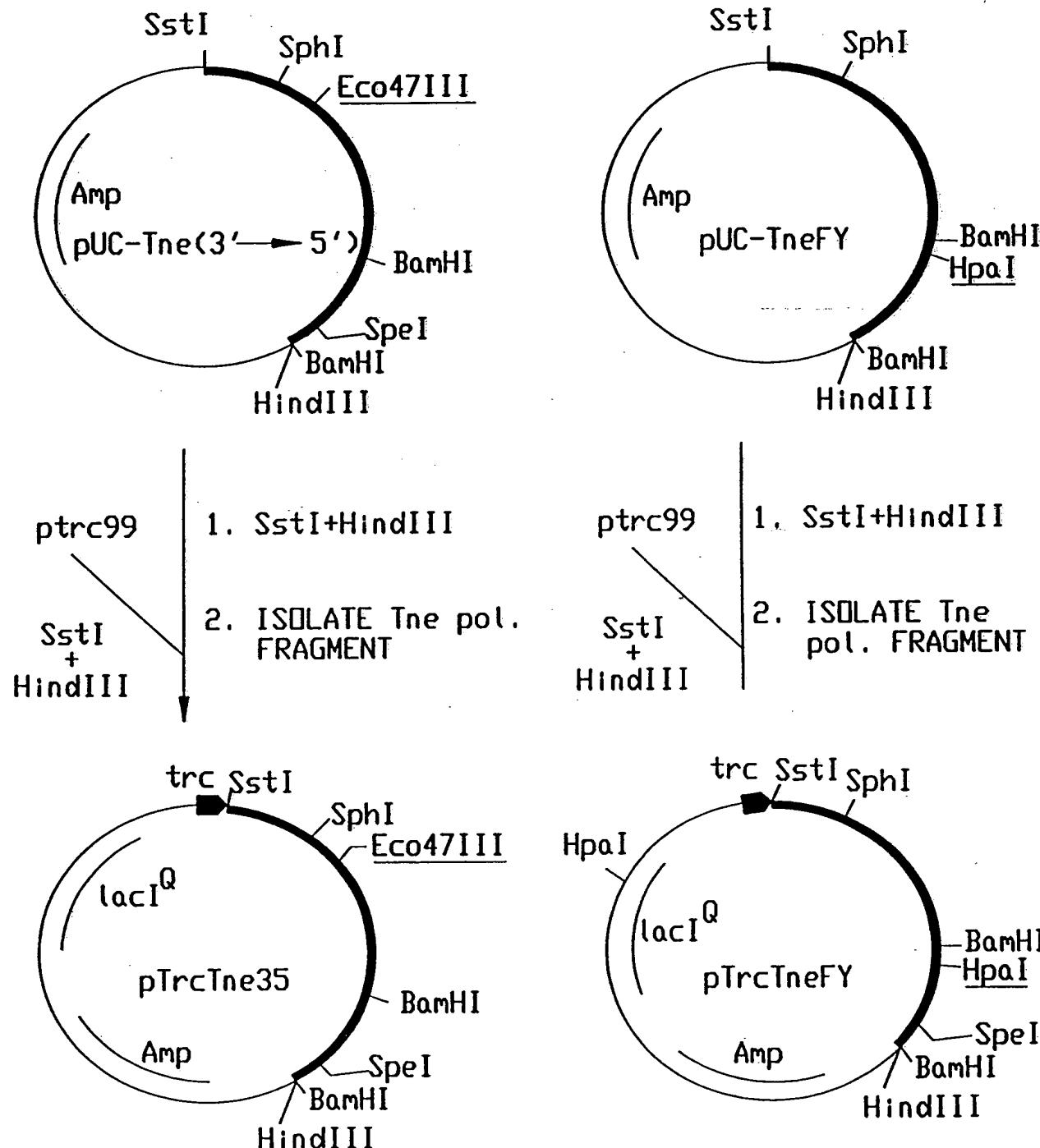


FIG.6B

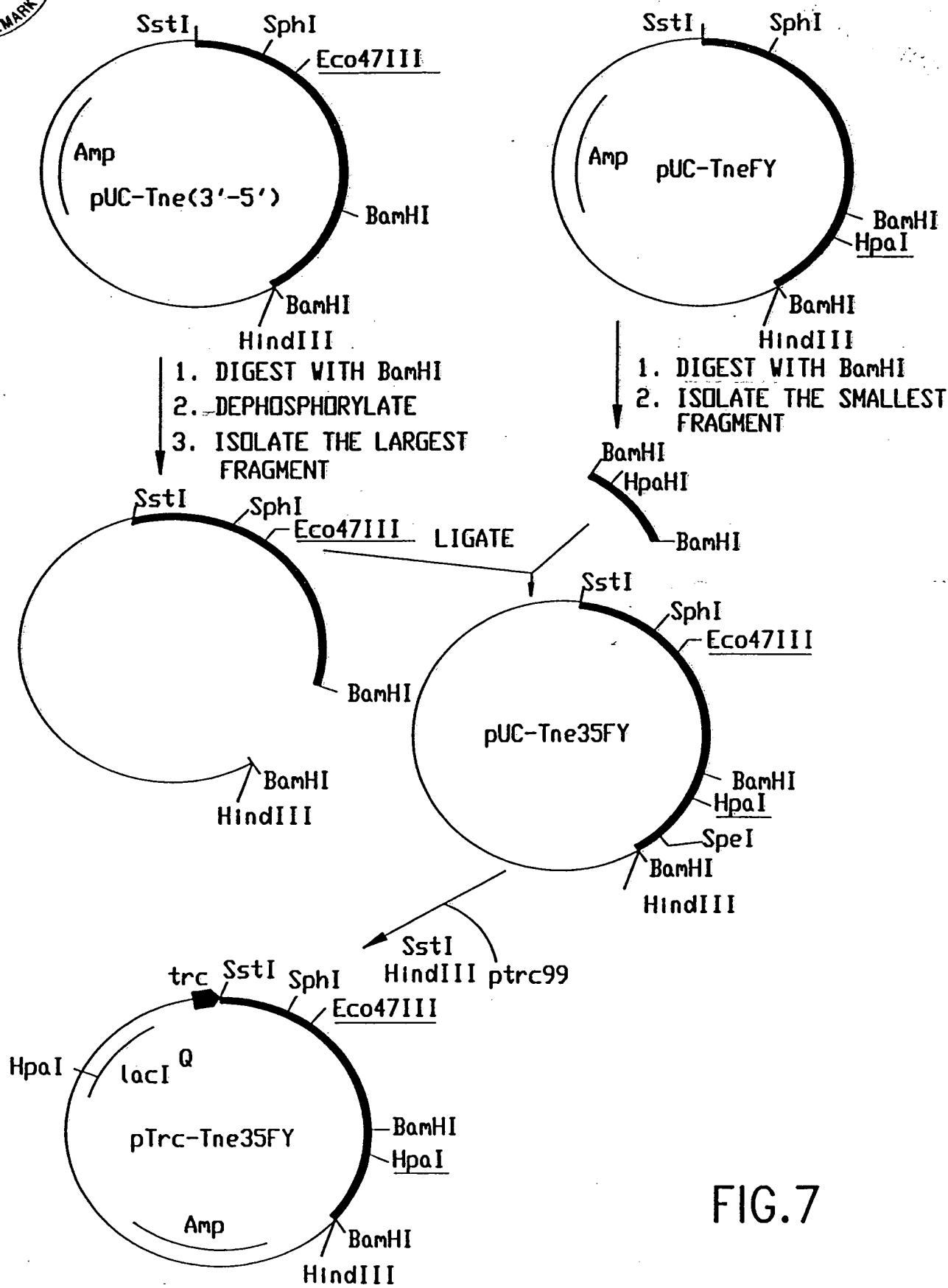


FIG.7

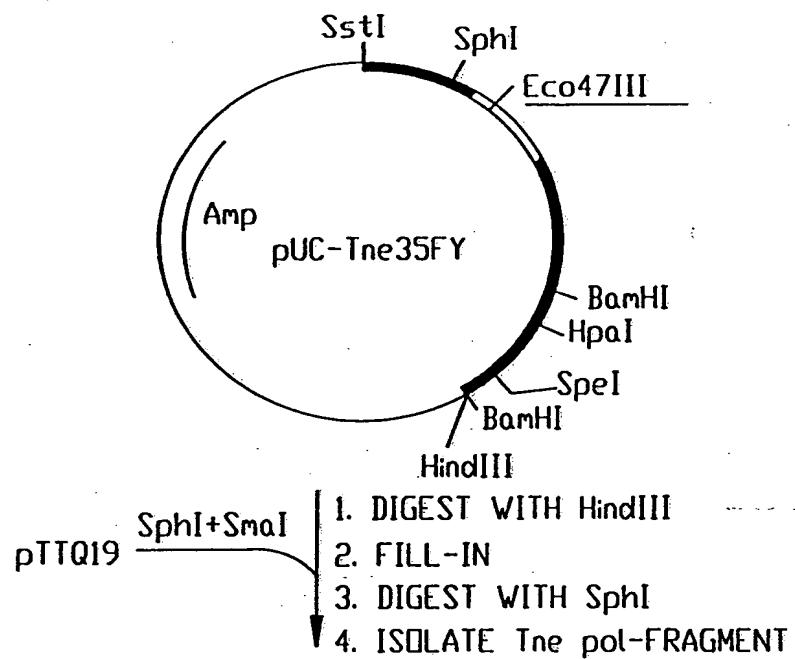
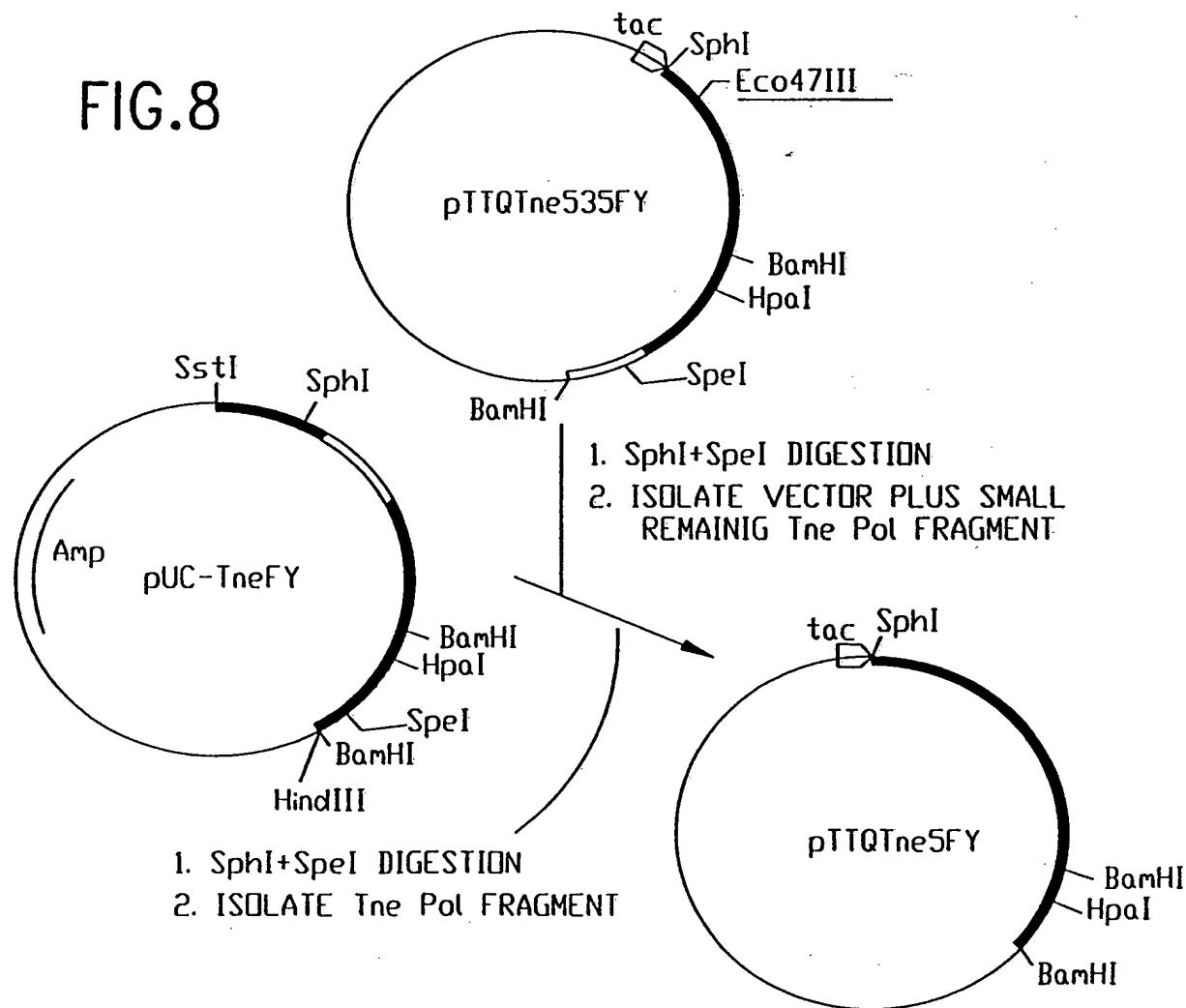


FIG.8





A                    B  
ACGT      GATC



FIG. 9



FIG.10

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A      B      C

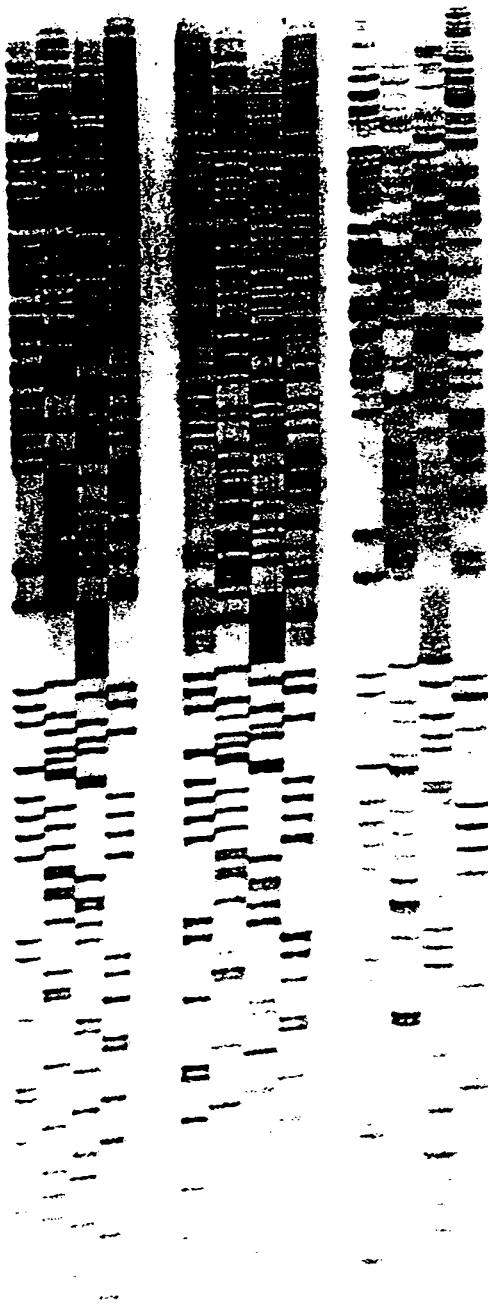


FIG.11

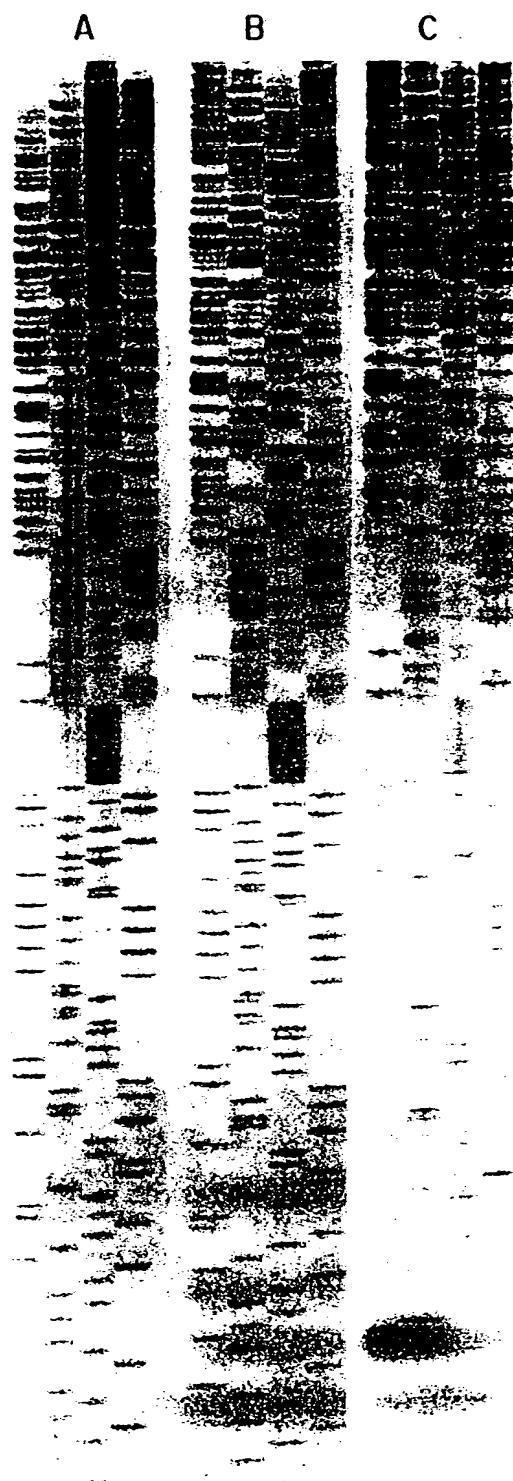


FIG.12

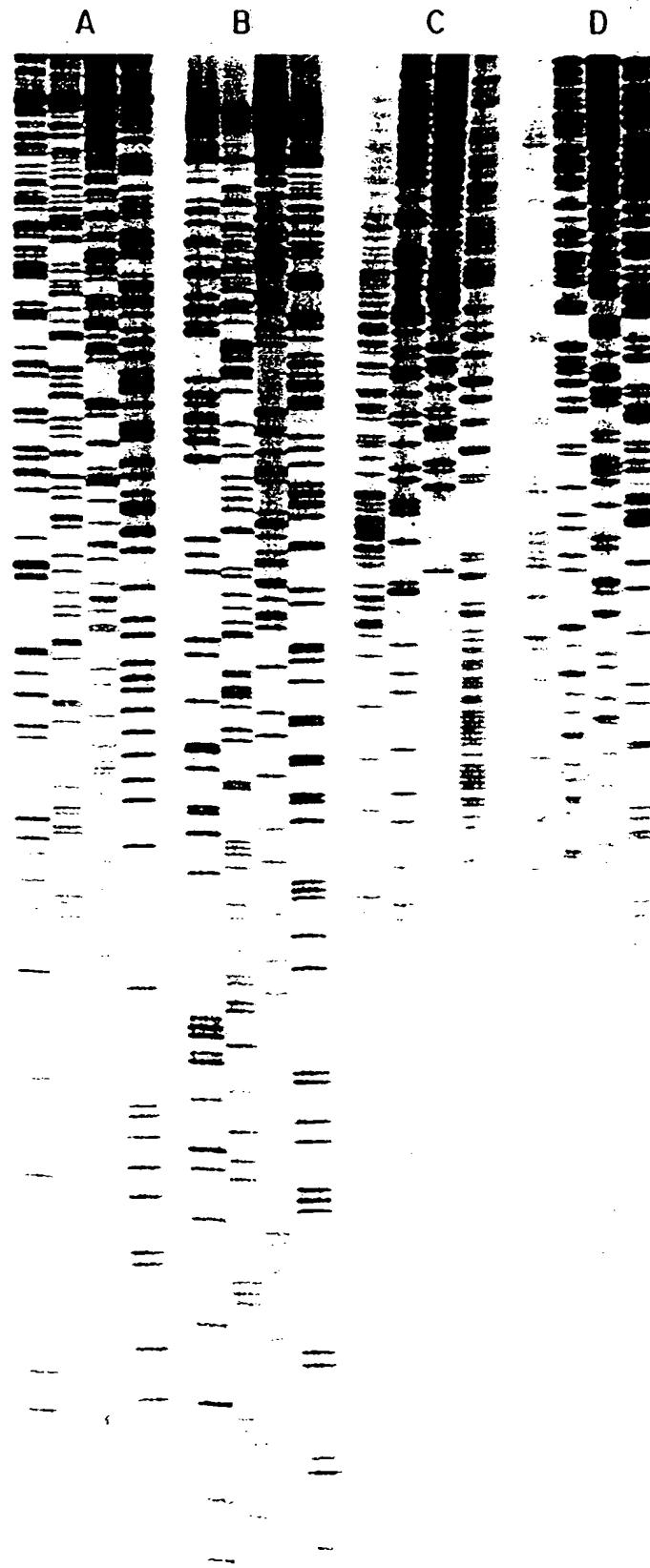


FIG. 13

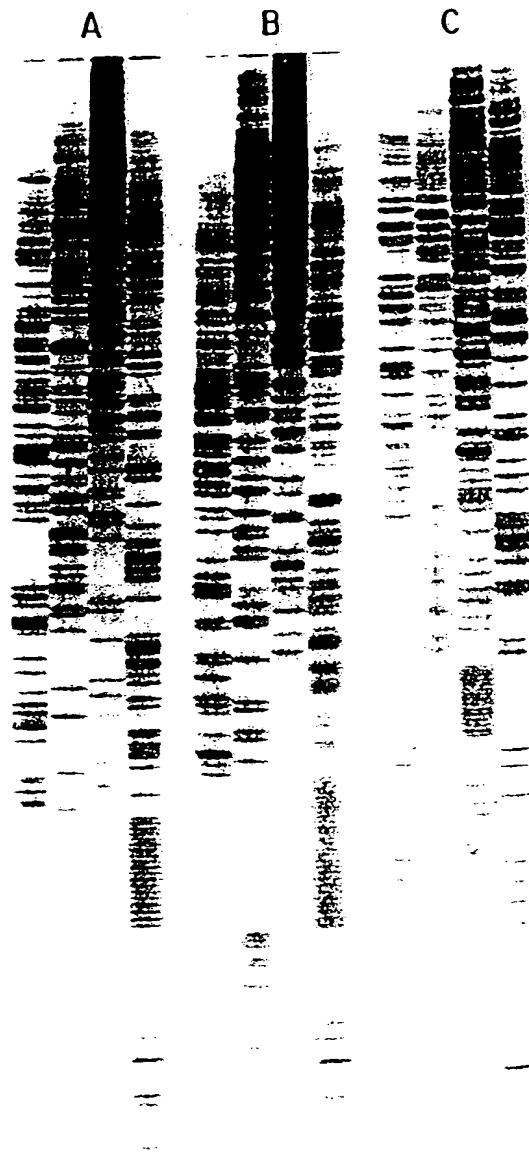


FIG.14A

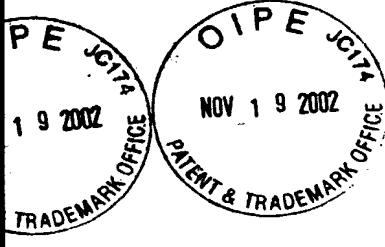


FIG. 14B

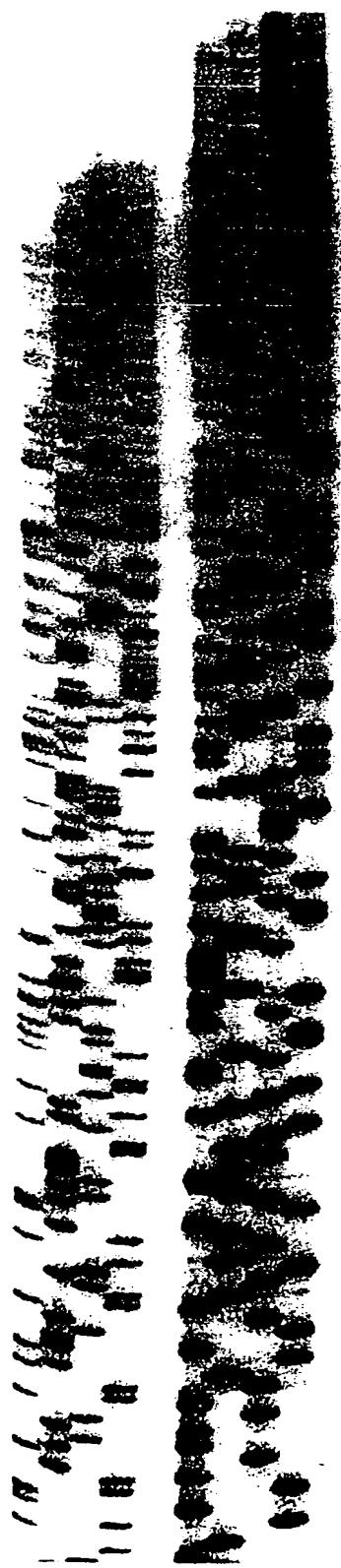


FIG. 15

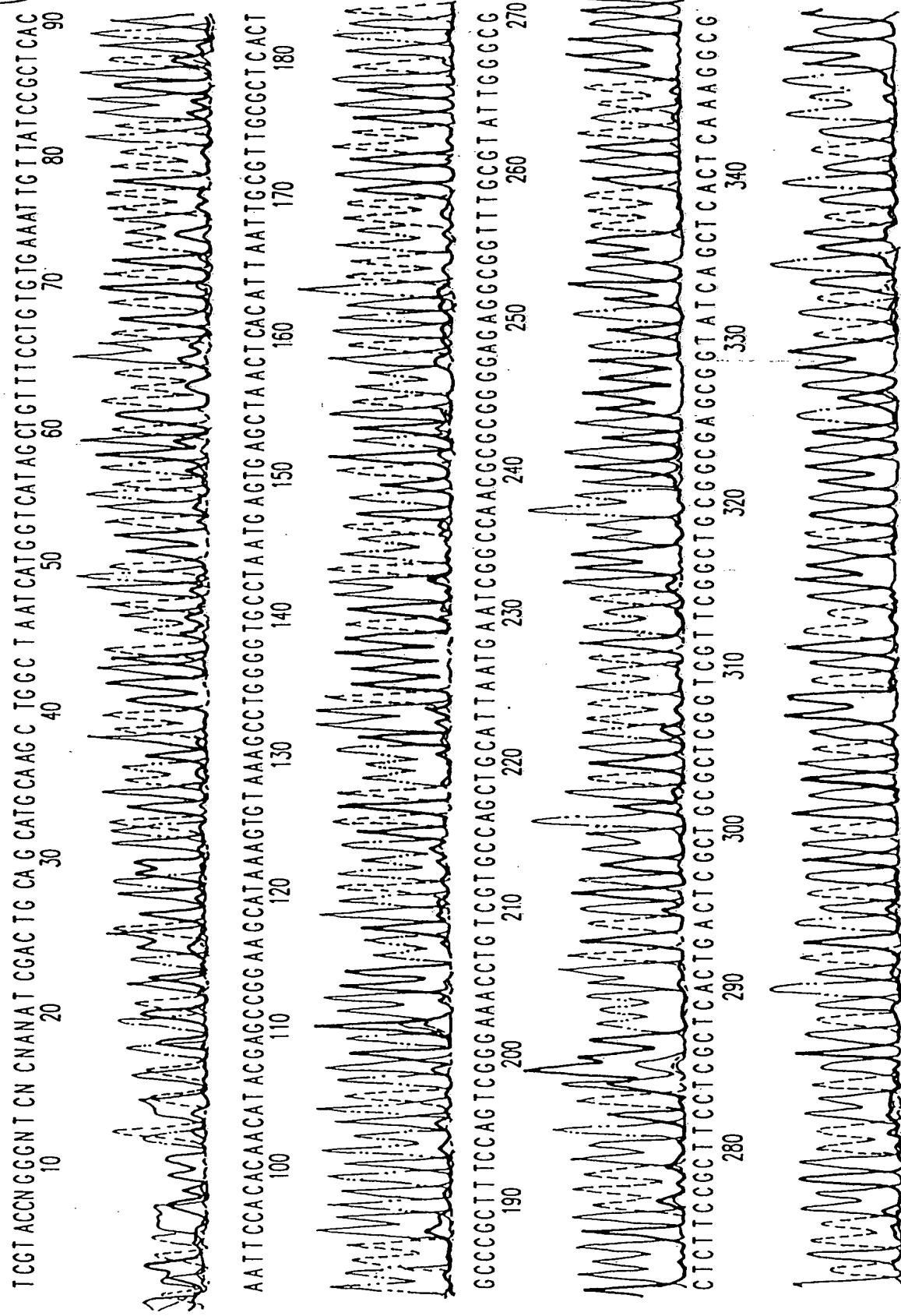


FIG. 16A

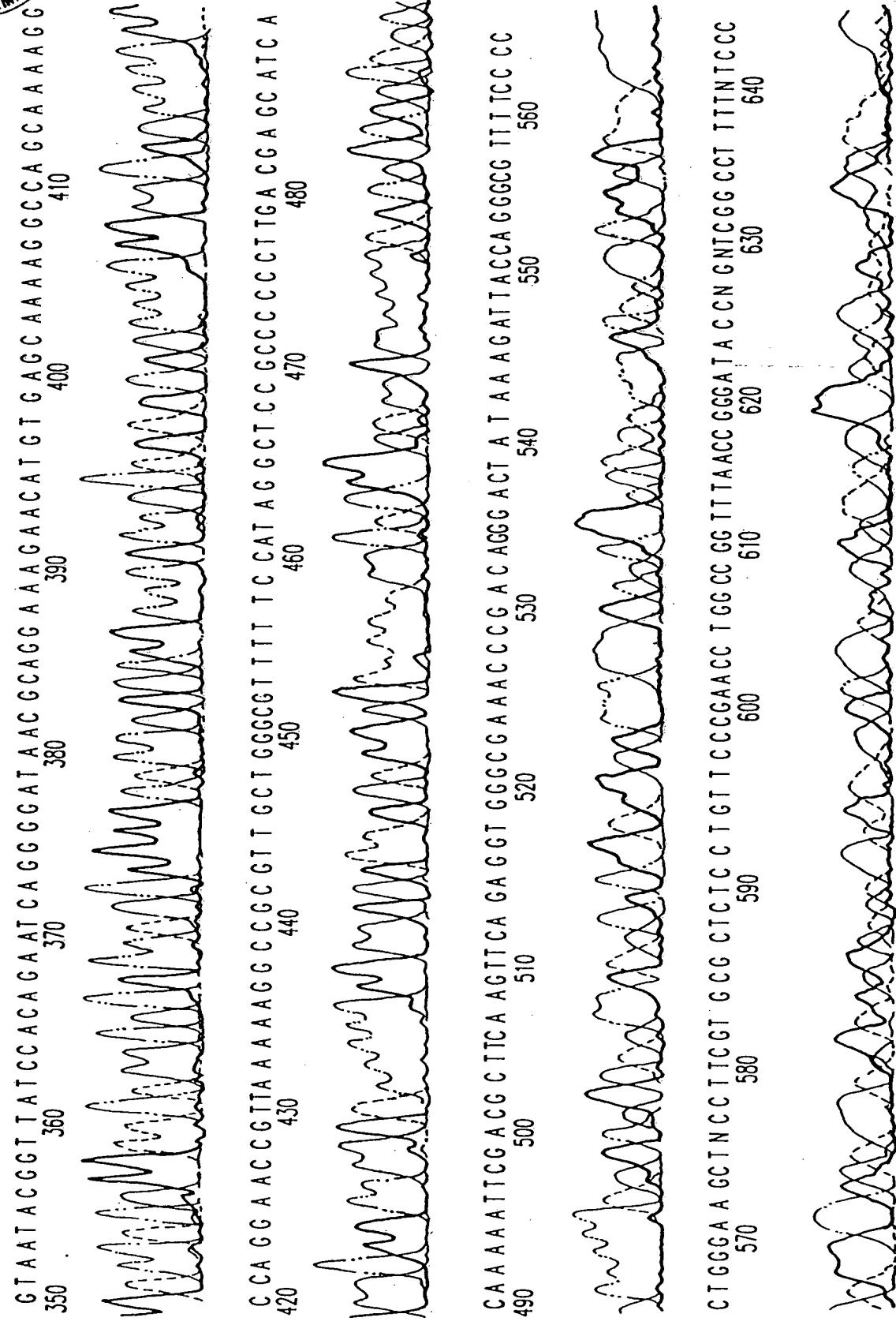


FIG. 16B



C T T N G G G G G A A N C C T T G C C G N T T T T I N G N A A A N G C T A A G G C T T  
650 660 670 680



FIG. 16C

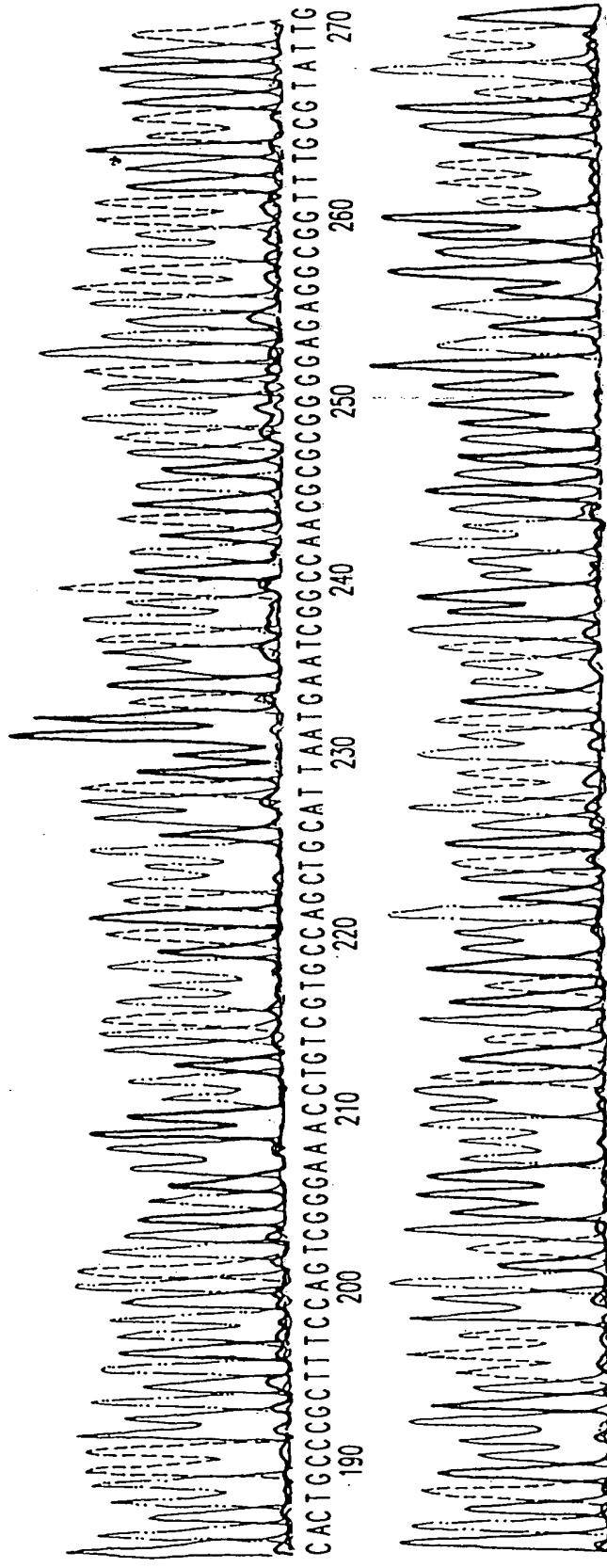
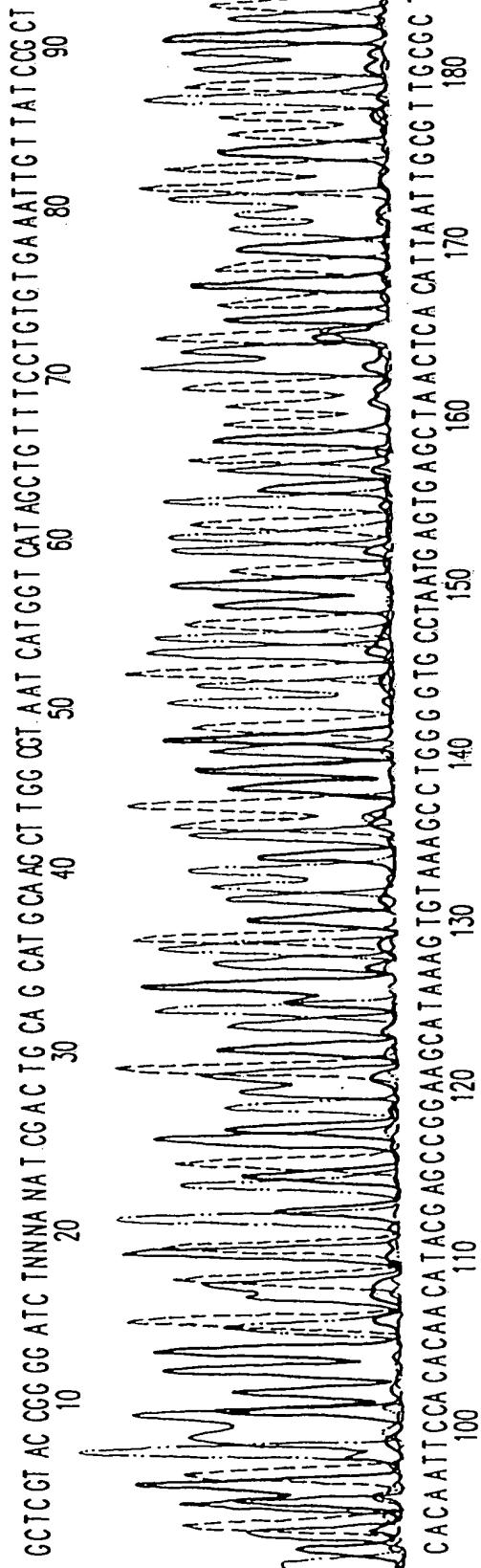
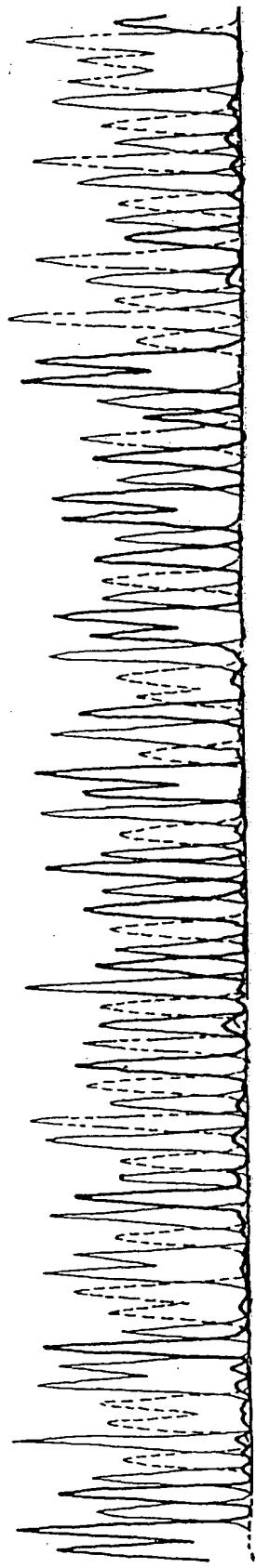


FIG. 16D



GGCGCTCTTCCGCTTCCTGACTCCCTGCCTTCGGCTGCCGAGCGGCTGCTACCTCAACTAAC  
280 290 300 310 320 330 340 350



GCGGTAAATACGGTTATCCACAGAATCAGGGATAACGCCAGGAAAGAACATGTGAGCAAAGCCAGCAA  
360 370 380 390 400 410 420

AGCCAGGAACCCGTAAGGGCGGCCCTTGCTGGCGTTTCTTCCATTAGGCTCCCTGANGACC  
430 440 450 460 470 480

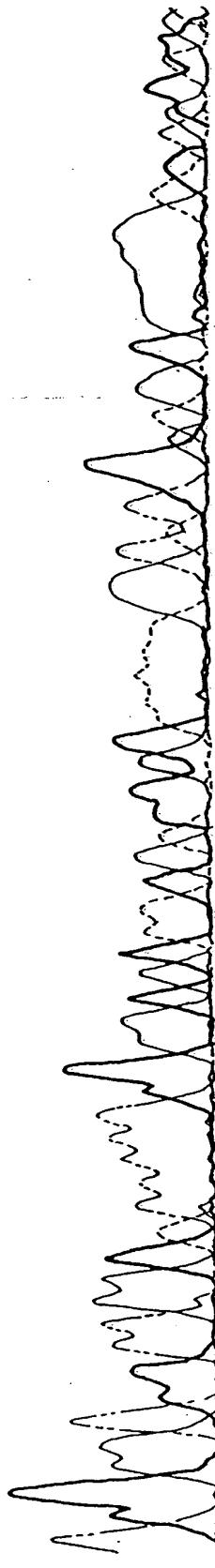


FIG. 16E



ATCANA A AAATCGA NGCT C A N G T C A N A G G T G G C G A AAC CC G AC A GG NC T AT TAAAG AT NCCAGG CGTTT  
490 500 510 520 530 540 550 560



CCCCC TGGG AAC GCT CCC T C G T C GGCCT C T C CT GG TT NGG NN CCCT GN CGG NT TA CGG QGGAT AAC TT GT TCCGG NT T NT  
570 580 590 600 610 620 630 640



CCCC T TNGGG AAAA NGGT GCGGG T TTTNT NAA AAC GCTCAA GGCTGGTANG  
650 660 670 680 690 700



FIG. 16F

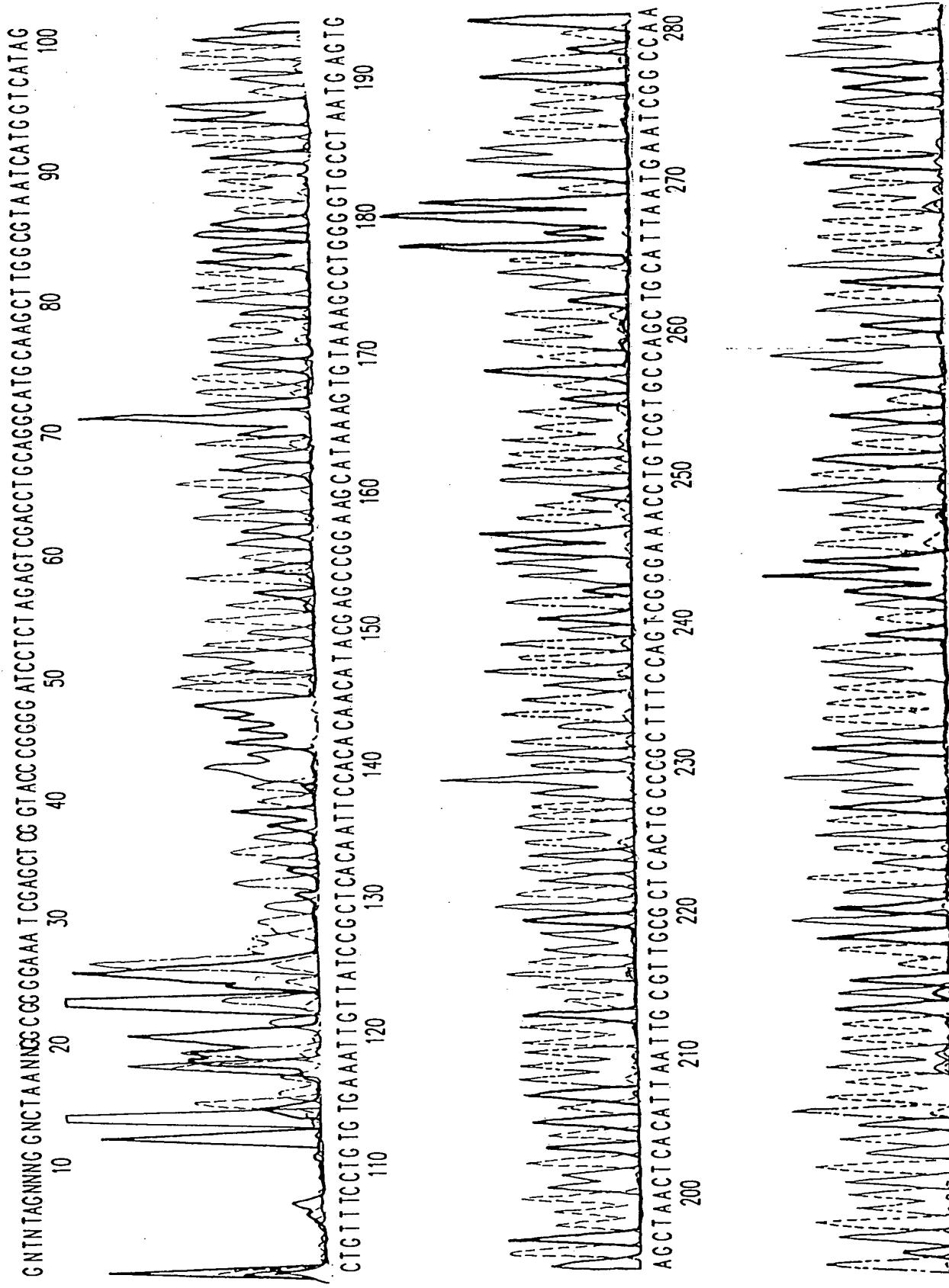
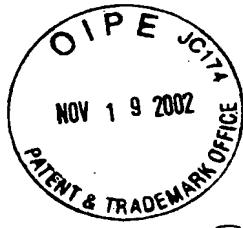
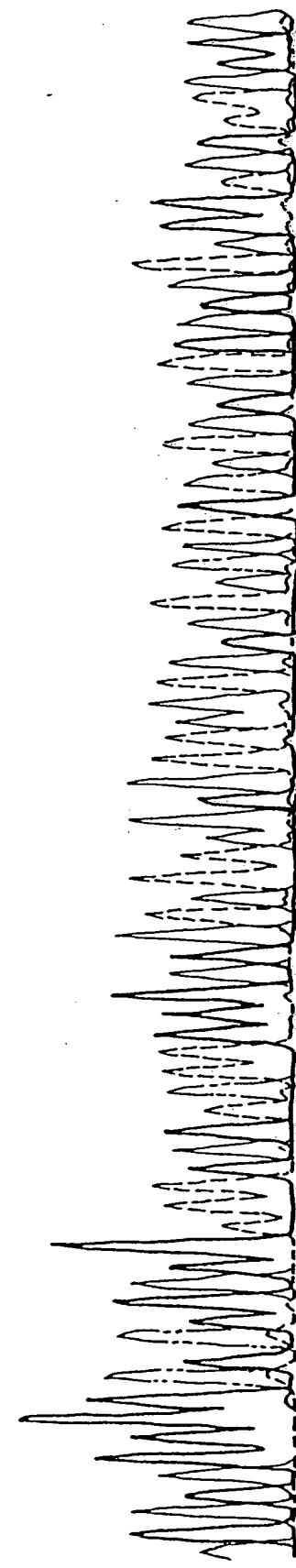


FIG. 17A



CGCG CGGGAGGGGTTTGCATTGGCGCTCTCCCTTCGGCTCACTGACTCGCTGGCTCCGTTCGCC  
290 300 310 320 330 340 350 360



TGGGGCGAGGGTATCAGCTCACTCAAAAGGGGGTAATACGGTTATCCACAGAAAATCAGGGGATAACGCCAGG  
370 380 390 400 410 420 430

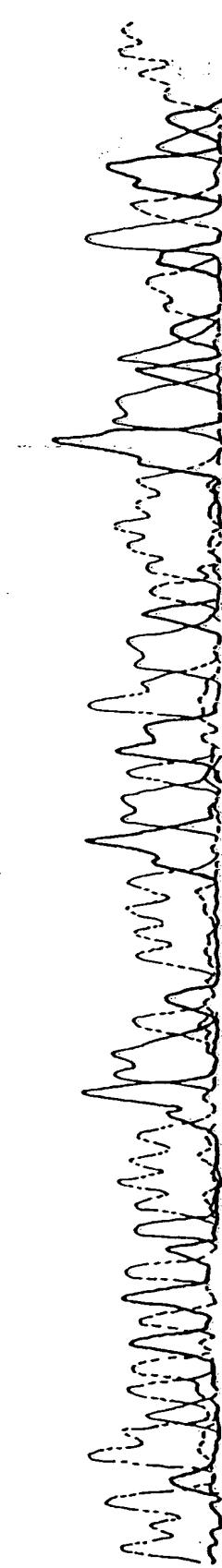
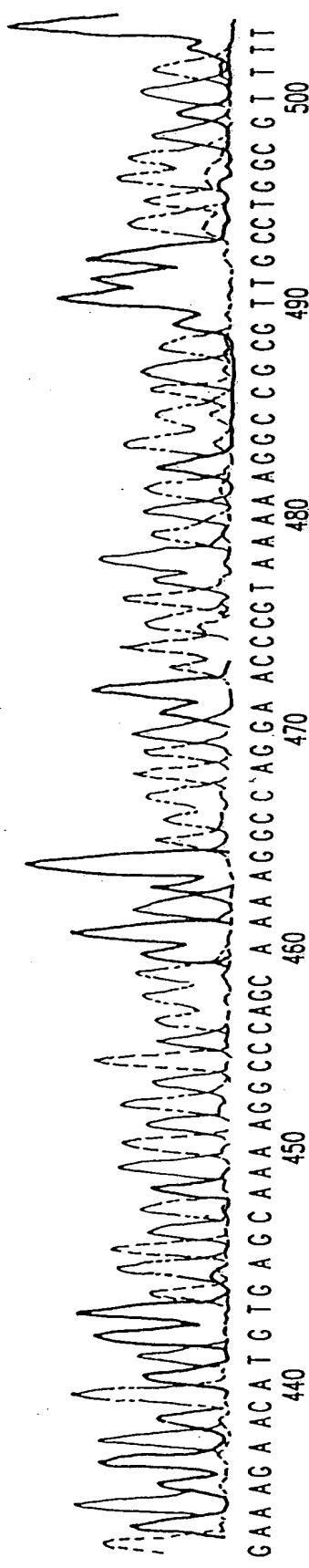
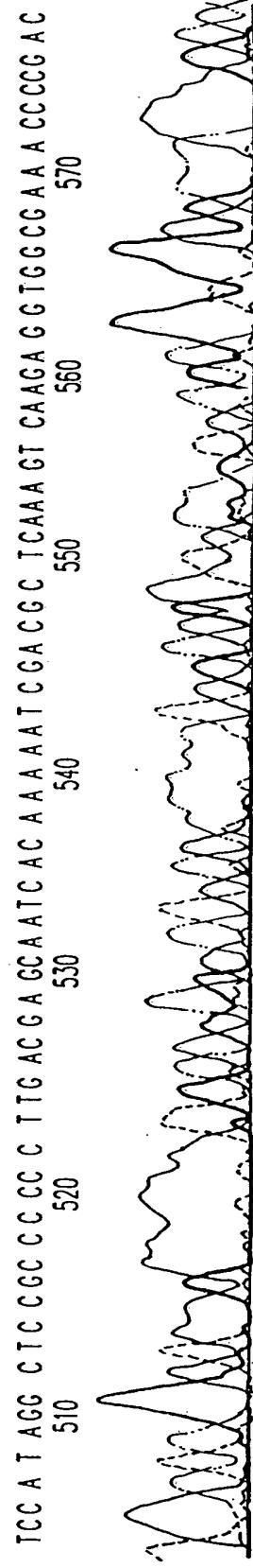


FIG. 17B



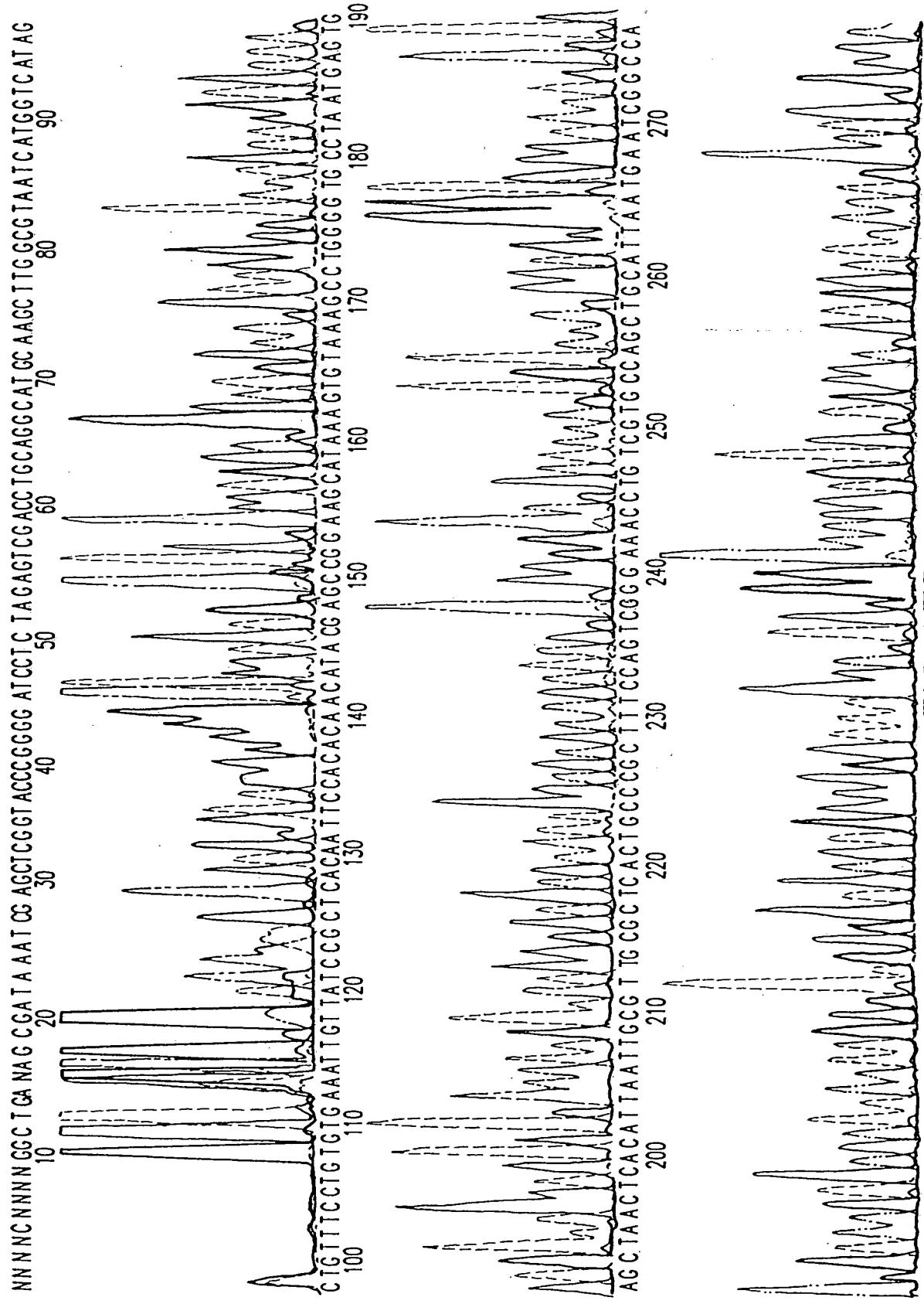


FIG. 17D



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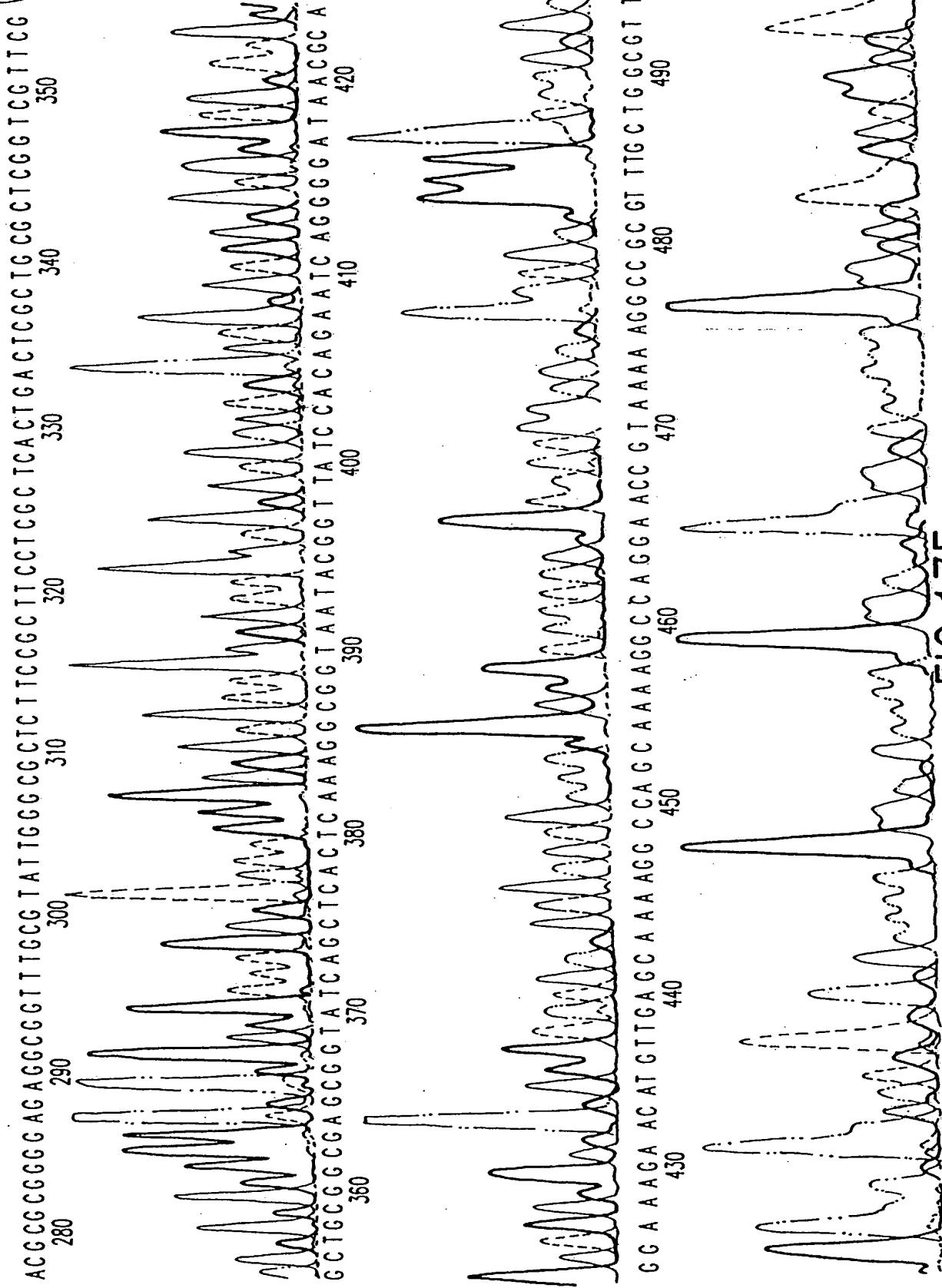


FIG. 17E



TTTCCCATAGGCCTCGCCCCCTTGACGAAATCGACGGCTCAATTAGAAGTTGGCGAAAACTGG  
500 510 520 530 540 550 560

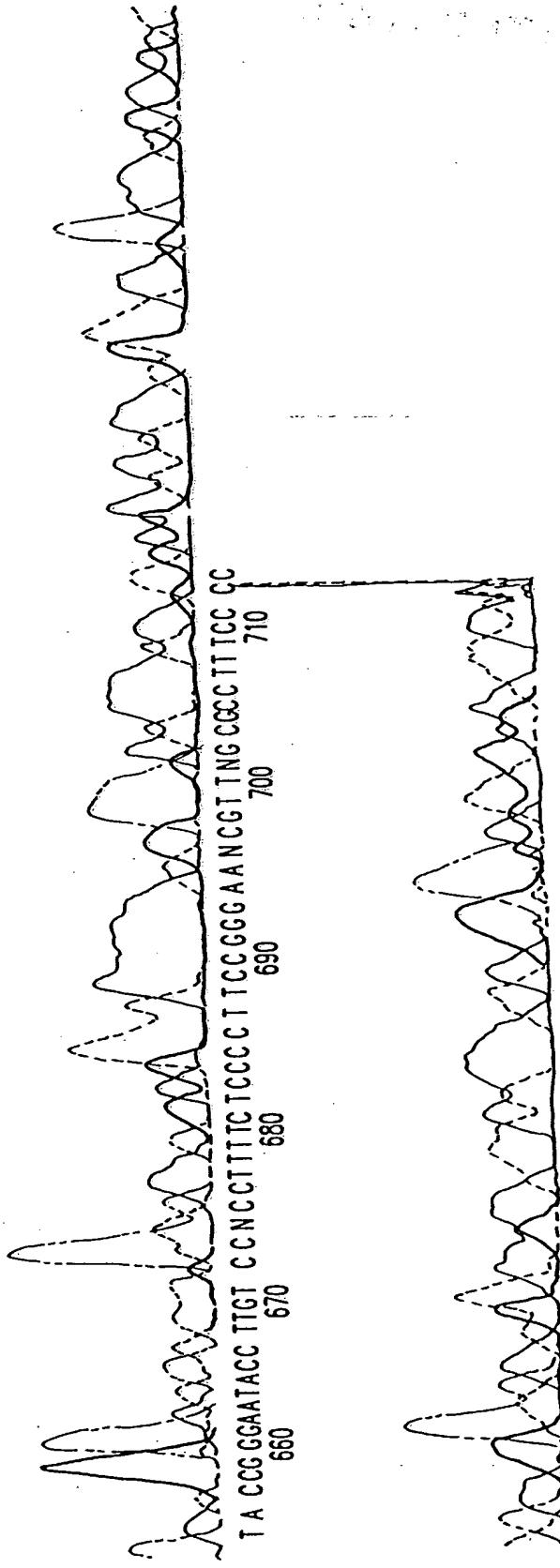
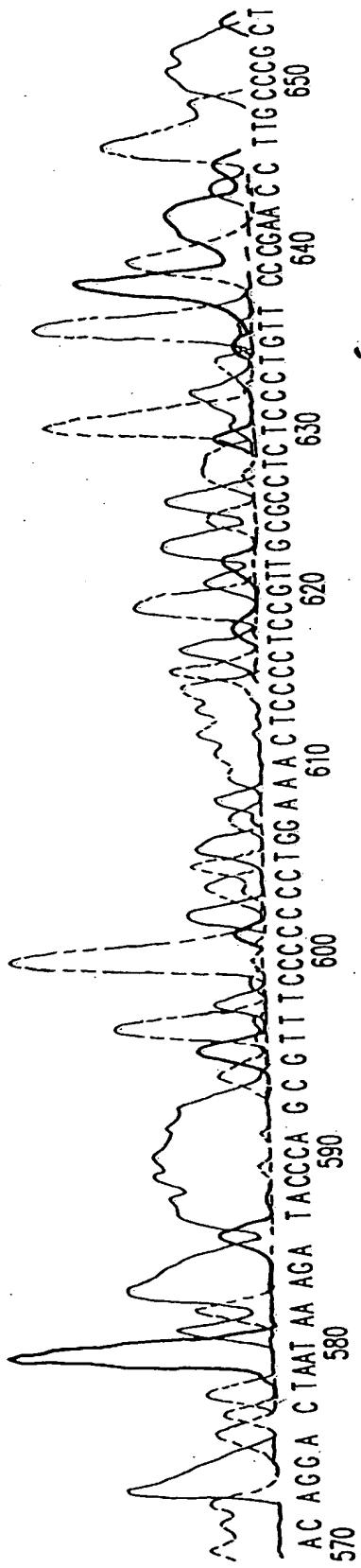


FIG. 17F



TTTCCCCATAGGCCTCGGCCCCCCTTGTACCAACCTTACAGAACCTGGCGAAAGTTGGCGAAAAACCCG  
500 510 520 530 540 550 560

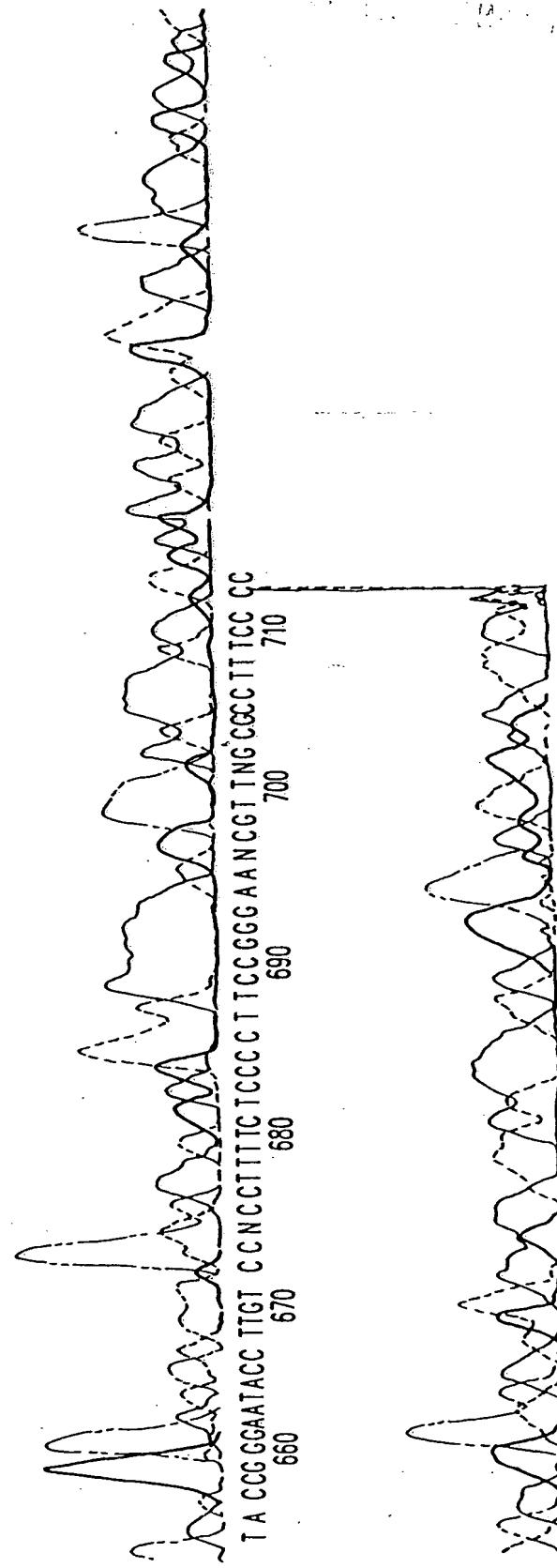
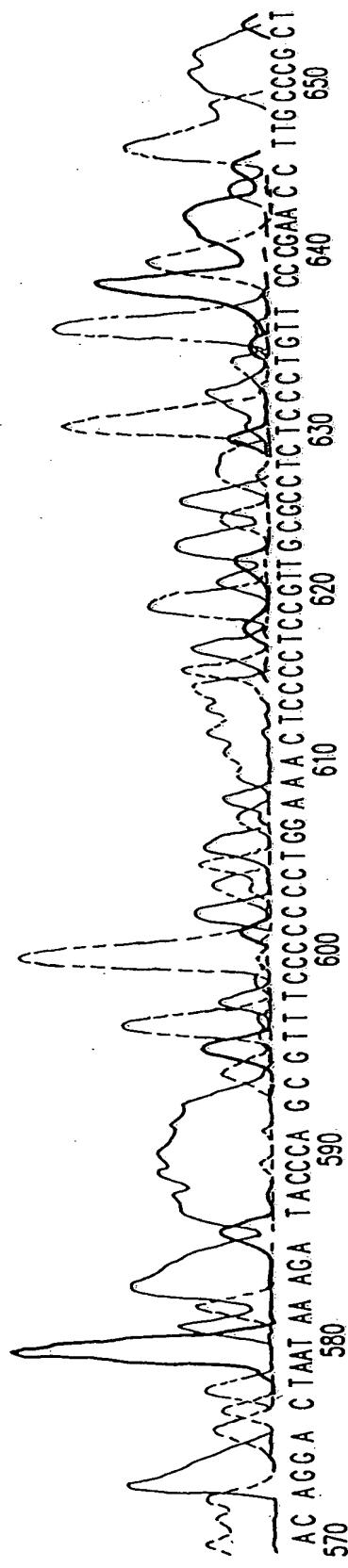


FIG. 17F